

Paolo Missier

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

115
papers

1,727
citations

19
h-index

38
g-index

137
ext. papers

2,076
ext. citations

2.2
avg, IF

4.53
L-index

#	Paper	IF	Citations
115	Knowledge-Driven Data Ecosystems Toward Data Transparency. <i>Journal of Data and Information Quality</i> , 2022 , 14, 1-12	2.5	4
114	The Right (Provenance) Hammer for the Job: A Comparison of Data Provenance Instrumentation. <i>Advanced Information and Knowledge Processing</i> , 2021 , 25-45	0.3	
113	A customisable pipeline for the semi-automated discovery of online activists and social campaigns on Twitter. <i>World Wide Web</i> , 2021 , 24, 1-37	2.9	0
112	Using Wearable Activity Trackers to Predict Type 2 Diabetes: Machine Learning-Based Cross-sectional Study of the UK Biobank Accelerometer Cohort. <i>JMIR Diabetes</i> , 2021 , 6, e23364	2.7	3
111	Machine learning in predicting respiratory failure in patients with COVID-19 pneumonia-Challenges, strengths, and opportunities in a global health emergency. <i>PLoS ONE</i> , 2020 , 15, e0239172	3.7	19
110	Capturing and querying fine-grained provenance of preprocessing pipelines in data science. <i>Proceedings of the VLDB Endowment</i> , 2020 , 14, 507-520	3.1	7
109	Abstracting PROV provenance graphs: A validity-preserving approach. <i>Future Generation Computer Systems</i> , 2020 , 111, 352-367	7.5	2
108	Why-Diff: Exploiting Provenance to Understand Outcome Differences From Non-Identical Reproduced Workflows. <i>IEEE Access</i> , 2019 , 7, 34973-34990	3.5	1
107	Increasing phenotypic annotation improves the diagnostic rate of exome sequencing in a rare neuromuscular disorder. <i>Human Mutation</i> , 2019 , 40, 1797-1812	4.7	9
106	Targeted therapies for congenital myasthenic syndromes: systematic review and steps towards a treataboloome. <i>Emerging Topics in Life Sciences</i> , 2019 , 3, 19-37	3.5	25
105	Toward trusted open data and services. <i>Internet Technology Letters</i> , 2019 , 2, e69	1.3	4
104	Report on the First International Workshop on Incremental Re-computation. <i>SIGMOD Record</i> , 2019 , 47, 35-38	1.1	
103	A Customisable Pipeline for Continuously Harvesting Socially-Minded Twitter Users. <i>Lecture Notes in Computer Science</i> , 2019 , 91-106	0.9	1
102	Toward a Decentralized, Trust-Less Marketplace for Brokered IoT Data Trading Using Blockchain 2019 ,		21
101	VazaDengue: An information system for preventing and combating mosquito-borne diseases with social networks. <i>Information Systems</i> , 2018 , 75, 26-42	2.7	17
100	Selective and Recurring Re-computation of Big Data Analytics Tasks: Insights from a Genomics Case Study. <i>Big Data Research</i> , 2018 , 13, 76-94	3.7	3
99	Analyzing Social Network Images with Deep Learning Models to Fight Zika Virus. <i>Lecture Notes in Computer Science</i> , 2018 , 605-610	0.9	3

98	Provenance Annotation and Analysis to Support Process Re-computation. <i>Lecture Notes in Computer Science</i> , 2018 , 3-15	0.9	
97	Leveraging Blockchain to Enable Smart-Health Applications 2018 ,		9
96	Versioned-PROV: A PROV Extension to Support Mutable Data Entities. <i>Lecture Notes in Computer Science</i> , 2018 , 87-100	0.9	2
95	TAPER: query-aware, partition-enhancement for large, heterogenous graphs. <i>Distributed and Parallel Databases</i> , 2017 , 35, 85-115	0.9	5
94	Mind my value 2017 ,		18
93	Recruiting from the Network: Discovering Twitter Users Who Can Help Combat Zika Epidemics. <i>Lecture Notes in Computer Science</i> , 2017 , 437-445	0.9	10
92	Preserving the Value of Large Scale Data Analytics over Time Through Selective Re-computation. <i>Lecture Notes in Computer Science</i> , 2017 , 65-77	0.9	1
91	Tracking Dengue Epidemics Using Twitter Content Classification and Topic Modelling. <i>Lecture Notes in Computer Science</i> , 2016 , 80-92	0.9	17
90	Provenance and data differencing for workflow reproducibility analysis. <i>Concurrency Computation Practice and Experience</i> , 2016 , 28, 995-1015	1.4	19
89	Scalable and efficient whole-exome data processing using workflows on the cloud. <i>Future Generation Computer Systems</i> , 2016 , 65, 153-168	7.5	8
88	Data trajectories: tracking reuse of published data for transitive credit attribution. <i>International Journal of Digital Curation</i> , 2016 , 11, 1-16	0.9	4
87	The Lifecycle of Provenance Metadata and Its Associated Challenges and Opportunities. <i>Springer Proceedings in Business and Economics</i> , 2016 , 127-137	0.2	3
86	DataONE: A Data Federation with Provenance Support. <i>Lecture Notes in Computer Science</i> , 2016 , 230-234	0.9	3
85	Analyzing Provenance Across Heterogeneous Provenance Graphs. <i>Lecture Notes in Computer Science</i> , 2016 , 57-70	0.9	5
84	Clustering provenance facilitating provenance exploration through data abstraction 2016 ,		5
83	ProvGen: Generating Synthetic PROV Graphs with Predictable Structure. <i>Lecture Notes in Computer Science</i> , 2015 , 16-27	0.9	7
82	Access control and view generation for provenance graphs. <i>Future Generation Computer Systems</i> , 2015 , 49, 8-27	7.5	7
81	2015 ,		23

80	YesWorkflow: A User-Oriented, Language-Independent Tool for Recovering Workflow Information from Scripts. <i>International Journal of Digital Curation</i> , 2015 , 10, 298-313	0.9	47
79	ProvAbs: Model, Policy, and Tooling for Abstracting PROV Graphs. <i>Lecture Notes in Computer Science</i> , 2015 , 3-15	0.9	8
78	Provenance Storage, Querying, and Visualization in PBase. <i>Lecture Notes in Computer Science</i> , 2015 , 239-241	0.9	1
77	Provenance-Based Searching and Ranking for Scientific Workflows. <i>Lecture Notes in Computer Science</i> , 2015 , 209-214	0.9	1
76	SVI: A Simple Single-Nucleotide Human Variant Interpretation Tool for Clinical Use. <i>Lecture Notes in Computer Science</i> , 2015 , 180-194	0.9	2
75	A Platform for Analysing Stream and Historic Data with Efficient and Scalable Design Patterns 2014 ,		2
74	From Scripted HPC-Based NGS Pipelines to Workflows on the Cloud 2014 ,		5
73	Tweet My Street: A Cross-Disciplinary Collaboration for the Analysis of Local Twitter Data. <i>Future Internet</i> , 2014 , 6, 378-396	3.3	14
72	Report from the second workshop on scalable workflow enactment engines and technology (SWEETU3). <i>SIGMOD Record</i> , 2014 , 42, 73-77	1.1	0
71	On strategies for budget-based online annotation in human activity recognition 2014 ,		3
70	Measuring the impact of cognitive distractions on driving performance using time series analysis 2014 ,		4
69	Distilling structure in Taverna scientific workflows: a refactoring approach. <i>BMC Bioinformatics</i> , 2014 , 15 Suppl 1, S12	3.6	6
68	The PBase Scientific Workflow Provenance Repository. <i>International Journal of Digital Curation</i> , 2014 , 9, 28-38	0.9	16
67	Measuring Population-Based Completeness for Single Nucleotide Polymorphism (SNP) Databases. <i>Studies in Computational Intelligence</i> , 2014 , 173-182	0.8	2
66	Forget Dimensions: Define Your Information Quality Using Quality View Patterns. <i>Synthese Library</i> , 2014 , 25-41	0.2	
65	Why linked data is not enough for scientists. <i>Future Generation Computer Systems</i> , 2013 , 29, 599-611	7.5	160
64	The W3C PROV family of specifications for modelling provenance metadata 2013 ,		73
63	Extracting PROV provenance traces from Wikipedia history pages 2013 ,		9

62	Report from the first workshop on scalable workflow enactment engines and technology (SWEETU2). <i>SIGMOD Record</i> , 2013 , 41, 60-64	1.1	
61	Measuring Data Completeness for Microbial Genomics Database. <i>Lecture Notes in Computer Science</i> , 2013 , 186-195	0.9	4
60	Reference Architectures to Measure Data Completeness across Integrated Databases. <i>Lecture Notes in Computer Science</i> , 2013 , 216-225	0.9	2
59	Predicting the Execution Time of Workflow Activities Based on Their Input Features 2012 ,		23
58	A PROV Encoding for Provenance Analysis Using Deductive Rules. <i>Lecture Notes in Computer Science</i> , 2012 , 67-81	0.9	6
57	Detecting Duplicate Records in Scientific Workflow Results. <i>Lecture Notes in Computer Science</i> , 2012 , 126-138	0.9	1
56	Modelling Provenance Using Structured Occurrence Networks. <i>Lecture Notes in Computer Science</i> , 2012 , 183-197	0.9	
55	. <i>IEEE Internet Computing</i> , 2011 , 15, 40-48	2.4	14
54	Simulating Taverna workflows using stochastic process algebras. <i>Concurrency Computation Practice and Experience</i> , 2011 , 23, 1920-1935	1.4	1
53	The Open Provenance Model core specification (v1.1). <i>Future Generation Computer Systems</i> , 2011 , 27, 743-756	7.5	396
52	Workflows to open provenance graphs, round-trip. <i>Future Generation Computer Systems</i> , 2011 , 27, 812-819	1.9	10
51	Achieving reproducibility by combining provenance with service and workflow versioning 2011 ,		14
50	The panel of experts cloud pattern 2011 ,		2
49	An experimental workflow development platform for historical document digitisation and analysis 2011 ,		5
48	eScience 2011 , 701-736		2
47	Workflows for Information Integration in the Life Sciences. <i>Lecture Notes in Computer Science</i> , 2011 , 215-225	0.9	1
46	Linking multiple workflow provenance traces for interoperable collaborative science 2010 ,		20
45	Functional Units: Abstractions for Web Service Annotations 2010 ,		5

44	ERGOT: A Semantic-Based System for Service Discovery in Distributed Infrastructures 2010 ,		14
43	The Evolution of myExperiment 2010 ,		5
42	Why Linked Data is Not Enough for Scientists 2010 ,		31
41	A Comparison of Using Taverna and BPEL in Building Scientific Workflows: the case of caGrid. <i>Concurrency Computation Practice and Experience</i> , 2010 , 22, 1098-1117	1.4	5
40	A formal semantics for the Taverna 2 workflow model. <i>Journal of Computer and System Sciences</i> , 2010 , 76, 490-508	1	19
39	Fine-grained and efficient lineage querying of collection-based workflow provenance 2010 ,		43
38	Taverna, Reloaded. <i>Lecture Notes in Computer Science</i> , 2010 , 471-481	0.9	71
37	Janus: From Workflows to Semantic Provenance and Linked Open Data. <i>Lecture Notes in Computer Science</i> , 2010 , 129-141	0.9	31
36	Understanding Collaborative Studies through Interoperable Workflow Provenance. <i>Lecture Notes in Computer Science</i> , 2010 , 42-58	0.9	12
35	A Linked Data Approach to Sharing Workflows and Workflow Results. <i>Lecture Notes in Computer Science</i> , 2010 , 340-354	0.9	2
34	2009 ,		2
33	Time-completeness trade-offs in record linkage using adaptive query processing 2009 ,		6
32	The data playground: An intuitive workflow specification environment. <i>Future Generation Computer Systems</i> , 2009 , 25, 453-459	7.5	8
31	Incorporating Domain-Specific Information Quality Constraints into Database Queries. <i>Journal of Data and Information Quality</i> , 2009 , 1, 1-31	2.5	6
30	Building Scientific Workflow with Taverna and BPEL: A Comparative Study in caGrid. <i>Lecture Notes in Computer Science</i> , 2009 , 118-129	0.9	16
29	Medical image processing workflow support on the EGEE grid with taverna 2009 ,		8
28	Data Provenance in Scientific Workflows 2009 , 46-59		1
27	Brokering infrastructure for minimum cost data procurement based on quality-quantity models. <i>Decision Support Systems</i> , 2008 , 45, 95-109	5.6	7

26	An ontology-based approach to handling information quality in e-Science. <i>Concurrency Computation Practice and Experience</i> , 2008 , 20, 253-264	1.4	9
25	Information quality in proteomics. <i>Briefings in Bioinformatics</i> , 2008 , 9, 174-88	13.4	25
24	Exploiting Provenance to Make Sense of Automated Decisions in Scientific Workflows. <i>Lecture Notes in Computer Science</i> , 2008 , 174-185	0.9	2
23	Data Lineage Model for Taverna Workflows with Lightweight Annotation Requirements. <i>Lecture Notes in Computer Science</i> , 2008 , 17-30	0.9	32
22	Grid metadata management: Requirements and architecture 2007 ,		3
21	Managing information quality in e-science 2007 ,		9
20	Requirements and Services for Metadata Management. <i>IEEE Internet Computing</i> , 2007 , 11, 17-25	2.4	12
19	Taverna Workflows: Syntax and Semantics 2007 ,		46
18	Architectural Patterns for the Semantic Grid 2007 , 119-134		
17	Accelerating Disease Gene Identification Through Integrated SNP Data Analysis 2007 , 215-230		1
16	Metadata Management in S-OGSA. <i>Lecture Notes in Computer Science</i> , 2007 , 712-719	0.9	
15	Semantic Support for Meta-Scheduling in Grids 2007 , 169-183		4
14	Managing Information Quality in e-Science Using Semantic Web Technology. <i>Lecture Notes in Computer Science</i> , 2006 , 472-486	0.9	7
13	Towards the Management of Information Quality in Proteomics 2006 ,		2
12	An overview of S-OGSA: A Reference Semantic Grid Architecture. <i>Web Semantics</i> , 2006 , 4, 102-115	2.9	50
11	Clustering Web pages based on their structure. <i>Data and Knowledge Engineering</i> , 2005 , 54, 279-299	1.5	40
10	Provider issues in quality-constrained data provisioning 2005 ,		1
9	Managing Information Quality in e-Science: A Case Study in Proteomics. <i>Lecture Notes in Computer Science</i> , 2005 , 423-432	0.9	4

8	Fine-grain web site structure discovery 2003 ,		7
7	Improving Data Quality in Practice: A Case Study in the Italian Public Administration. <i>Distributed and Parallel Databases</i> , 2003 , 13, 135-160	0.9	13
6	Electronic Government. <i>Lecture Notes in Computer Science</i> , 2003 ,	0.9	21
5	The Service to Businesses Project: Improving Government-to-Business Relationships in Italy. <i>Lecture Notes in Computer Science</i> , 2003 , 468-471	0.9	2
4	CitiTime: a system for rapid creation of portable next-generation telephony services. <i>Computer Networks</i> , 2001 , 35, 579-595	5.4	1
3	Java call control, coordination, and transactions 2000 , 38, 108-114		9
2	A framework for analyzing virtual enterprise infrastructure 1999 ,		4
1	An Overview of S-OGSA: A Reference Semantic Grid Architecture. <i>SSRN Electronic Journal</i> ,	1	1